

Notice of References Cited	Application/Control No. 09/806,178	Applicant(s)/Patent Under Reexamination NAGAI ET AL.	
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U.S. PATENT DOCUMENTS

*		Document Number Country Code-Number-Kind Code	Date MM-YYYY	Name	Classification
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NON-PATENT DOCUMENTS

*		Include as applicable: Author, Title Date, Publisher, Edition or Volume, Pertinent Pages)
✓	U	Bork (2000) "Powers and Pitfalls in Sequence Analysis: The 70% Hurdle." Genome Research 10:398-400
✓	V	Skolnick and Fetrow (2000) "From gene to protein structure and function: novel applications of computational approaches in th genomic era." Trends in Biotech. 18(1): 34-39
✓	W	Doerks et al., (June 1998) "Protein annotation: detective work for function prediction." Trends in Genetics 14(6): 248-250
✓	X	Smith and Zhang (November 1997) "The challenges of genome sequence annotation or 'The devil is in the details'." Nature Biotechnology 15:1222-1223

*A copy of this reference is not being furnished with this Office action. (See MPEP § 707.05(a).)
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<input checked="" type="checkbox"/>	V	Bork and Bairoch (October 1996) "Go hunting in sequence databases but watch out for the traps." Trends in Genetics 12(10): 425-427
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